1/11

Structure of heme

FIGURE 1

Alpha globin sequence:

2/11

							_		
4 3	06	135	180	225	270	315	360	405	
ට ල් ල්	GAG Glu	CAC	AAG Lys	GAC Asp	AAG Lys	CTG Leu	Ala Ala	GTG Val	
TGG Trp	CTG Lev	P 5	3 3 3	GAC Asp	CAC	TGC Cys	CCT Pro	ACC Thr	
Aa GC	AB 600	TTC Phe	EAC ES	GTG Val	Ala Ala	CAC	Agg	AGC Ser	
GCC Ala	GAG Glu	TAC Tyr	3 000 €	CAC	CAC	AGC Ser	TTC Phe	GTG Val	
AAG Lys	Ala A	AOC Thr	AAG Lys	603 Ala	CTG Leu	CTA Leu	6 46	TCT Ser	
GTC Val	GGT Gly	AAG Lys	GTT Val	GTG Val	GAC Asp	CTC	Aa SS	GCT Ala	
AAC Asn	TAT Tyr	ACC Thr	S S	Ala Ala	AGC	AAG Lys	8 g	CTG Leu	
ACC	8	AOC Thr	Aa Aa	AAC Asn	CTG Leu	TC Phe	CTC	Phe	6
AAG Lys	8 8 8	8 g	TCT Ser	A00 Thr	Aa Aa	AAC	CAC	AAG Lys	, and a second
GAC	GCT Ala	TC Phe	3 §	CTG Leu	TCC Ser	GTC Val	8 8 8	Asp	Arg
S Ala Ala	CAC His	TCC Ser	CA Els	8 8 8 8	CTG Leu	8 9 9 5	Ala Ala	CTG Lev	TAC Tyr
CCT Pro	8 80	CTG Lee	AGC Ser	GAC Asp	GOG Ala	GAC ASP	CTG Lee	Ser	AAA Lys
TCT Ser	3 1 1 1 1 1 1 1 1	TT Safe	CTG Lee	8 8 8	AAC Asn	GTG Val	ACC Thr	8 8 8 8	TCC Ser
CTG Lev	CTT Val	ATG Met	GAC. Asp	GTG Val	8 5	000 Arg	GTG Val	CAC	AQC Thr
GTG Val	AAG Lys	AGG Arg	11C	AAG Lys	ATG Met	CTT Lea	CTG Leu	GTG Val	CTG
	46	6	136	181	226	27.1	316	361	408

FIGURE 2

Beta globin sequence:

405 315 225 135 180 45 A&A Lys GCT Ala Cic Lee CTG Lee **8**€ GTG Val 17 g Arg Arg GTG Val **8** 8 € AGG Arg AGT Ser AAG Lys Cdg Leu TCC Ser 8 8 8 8 TTC Phe TTT GTG Val CTG Leu **8**89 **CC P c c** GAT Asp CTG SG e∦ A&A Lys AAC Asn CAC AAC Asn AGT Ser ACA TE TTT Phe ACT Thr 8 8 **8** 5 **8** 8 **8**8 CAT Phe Phe TC Pie **8** € **8**8 € Cal TAT Tyr TAT Tyr Şa Şa CCT Pro ATG Met \$ **₽** TTT 88 द्धार ब्रम् Arg \$ 88 Se eg ¥ ₹ CTG Lee GAT Asp ල් මේ ප් <u>G</u> GTT Val TCT द्धार अर् Sec Esc GTG Val **8** § GTG Val GCT Ala CTC Lee ¥ ₹ GCT Ala CT Val Lys **S S S** CAC **TGT** Cys AAG Lys **TGG Trp** GTG Val **8** 8 € GAT Asp **9** 8 E d GTG Val GTC Val CTC Lee CTG AAA Lys CCT Pro CCT Pro **8** 8 GAT Asp § 88 8 Pro Pro AAC Asn AAG Lys CTG AAG Lys ACT Thr TAC GTG Val Pro Pro AAT Asn GAC Asp GAC Asp GTG Val TCC Ser **3**€ **3**€ GTC Val AAC Asn ACT The T AAC Asn 1 A E TGT Cys CTG Lee CdG Lec CAT His Cata Val Cal Val Cd Lee HG Fe CAC H≥ SC ල ල් CAC His CTG Lee GAT Asp AAG Lys 8 8 CTG Lee CTG Lee AAG Lys GCT Ala **8**89 € 8 8 8 8 Cd Lee 406 361 226 271 136 181 46 91

FIGHRE 3

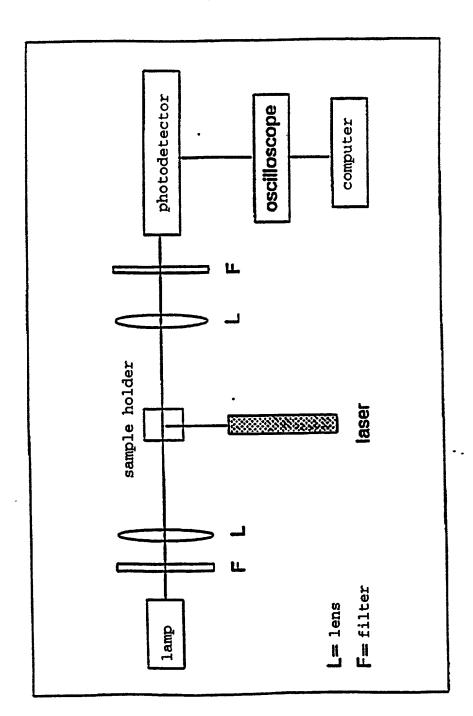
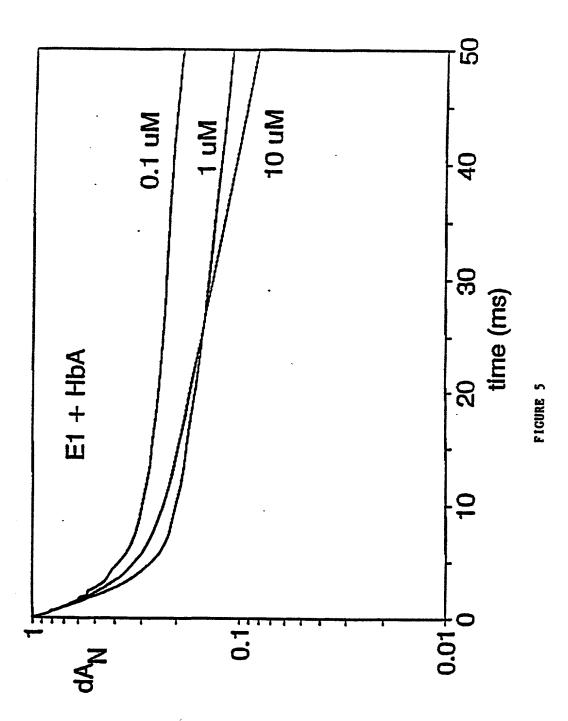
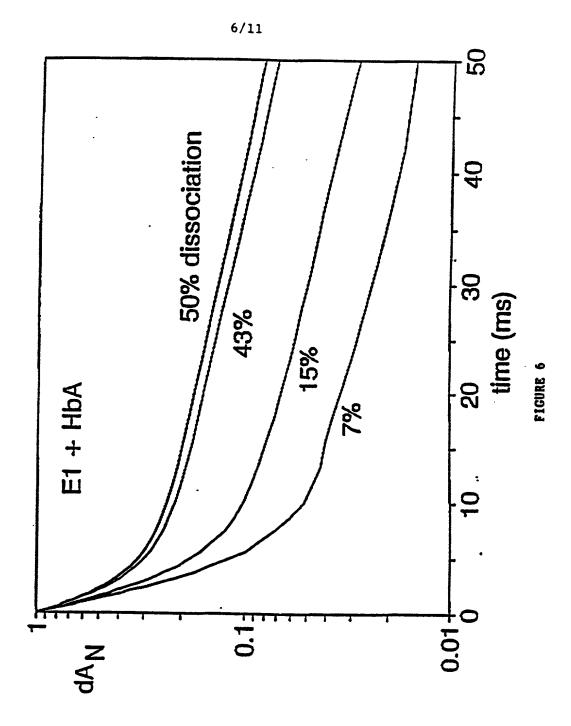


FIGURE 4





7/11

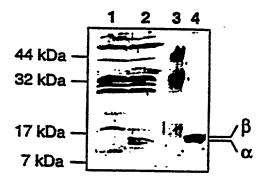


FIGURE 7

8/11

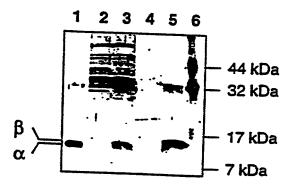


FIGURE 8

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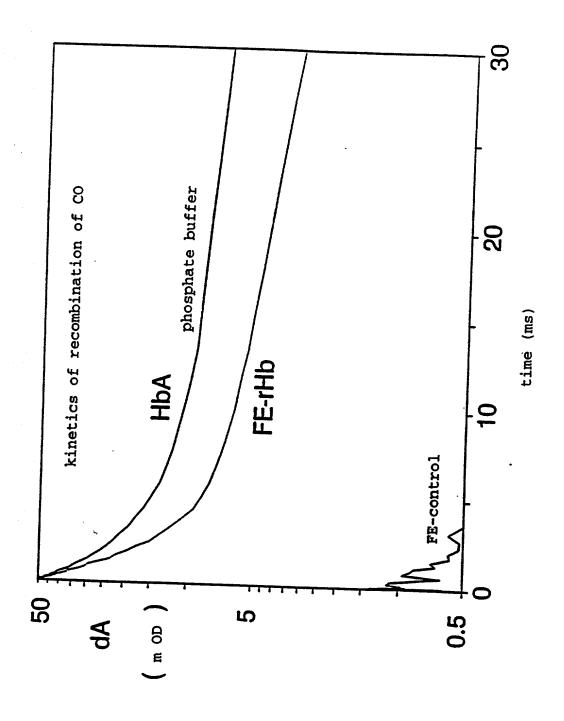


FIGURE 9

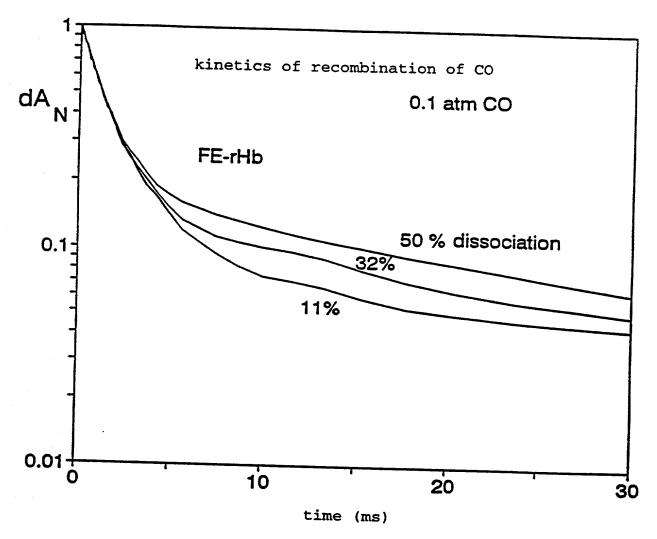


FIGURE 10

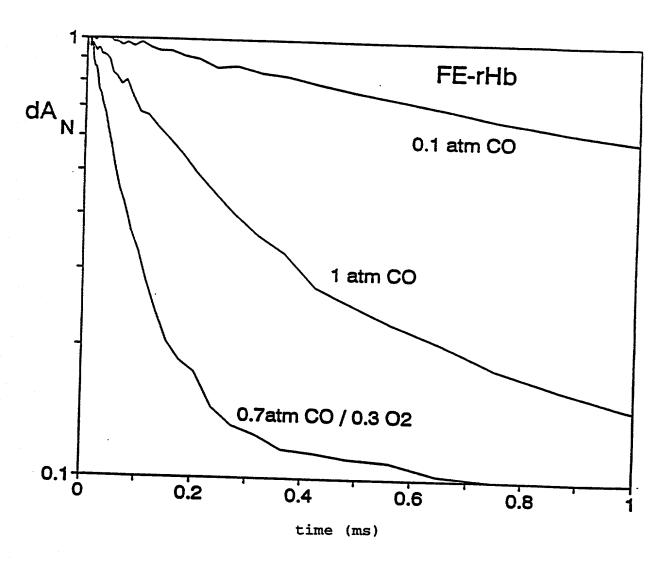


FIGURE 11